1 tgggtgt <u>etcccttgctctgccaacg</u> ttgttgattgttttc <u>atg</u> acattaatctacgtgc  Met Thr Leu Ile Tyr Val
61 cttcaatatttacaatggtcccctcaatcacacggattgtactggttaacattctgttgg 7 Pro Scr IIc Phe Thr Met Val Pro Ser IIe Thr Arg IIe Val Leu Val Asn IIe Leu Leu
121 cgacgttggttttgggagctgcagtccttccacgagacaacagaactgtttgcgggagtc 27 Ala Thr Leu Val Leu Gly Ala Ala Val Leu Pro Arg Asp Asp Arg Thr Val Cys Gly Ser
181 aactetgeacatggtggcacgacteeggegagataaacaceggtacteetgtacaggcag 47 Gln Leu Cys Thr Trp Trp His Asp Ser Gly Glu Ile Asn Thr Gly Thr Pro Val Gln Ala
241 gaaacgttcgacaatcccgaaagtactctgtccatgtgagcctggcagaccgtaaccaat 67 Gly Asn Val Arg Gln Ser Arg Lys Tyr Ser Val His Val Ser Leu Ala Asp Arg Asn Gln
301 tctacgactctttcgtatatgaatcgatacctaggaacggcaatggcagaatttattctc 87 Phe Tyr Asp Sor Phe Val Tyr Glu Scr IIc Pro Arg Asn Gly Asn Gly Arg IIe Tyr Ser
361 ccaccgacccacctaacagcaatacattgaatagtagcattgacgacggtatatcaatcg 107 Pro Thr Asp Pro Pro Asn Ser Asn Thr Leu Asn Ser Ser Ile Asp Asp Gly Ile Ser Ile
421 aaccateteteggeateaacatggettggteecagttegaatatagaegagatgtegaea 127 Glu Pro Ser Leu Gly Ile Asn Met Ala Trp Ser Gln Phe Glu Tyr Arg Arg Asp Val Asp
481. ttaagattactacaatcgatggctcaatattggatggccctttggacattgttattcggc 147 lle Lys Ile Thr Thr Ile Asp Gly Ser Ile Leu Asp Gly Pro Leu Asp Ile Val Ile Arg
541 cgacttctgttaagtactcagtcaaaagatgtgtgggtgg
601 atgateceaatggtegaaaattetetgttgagttaaagagtgacetttaeagttaeetet 187 Tyr Asp Pro Asn Gly Arg Lys Phe Ser Val Glu Leu Lys Ser Asp Leu Tyr Ser Tyr Le
661 ccgacggttcgcaatatgtgacctctggagggagcgtggttggt
721 ccctggtgatctttgccagccctttcttgccacgggatatggttcctcatatgacaccac 227 Ala Leu Val Ile Phe Ala Ser Pro Phe Leu Pro Arg Asp Met Val Pro His Met Thr Pro
781 acgacacccagacaatgaagccgggcccaatcaataatggggactggggttcaaagccta 247 His Asp Thr Gln Thr Met Lys Pro Gly Pro Ile Asn Asn Gly Asp Trp Gly Ser Lys Pro
841 tactctacttcccgcctggcgtatactggatgaacgaggatacctctggtaaccccggga 267 Ile Leu Tyr Phe Pro Pro Gly Val Tyr Trp Met Asn Glu Asp Thr Ser Gly Asn Pro Gly
901 ageteggeteaaateatatgeggetggateeeaatacetaetgggteeatetageeeeag 287 Lys Leu Gly Ser Asn His Met Arg Leu Asp Pro Asn Thr Tyr Trp Val His Leu Ala Pro
961 gagcctatgtgaaaggagccattgagtatttcacgaagcaaaatttctatgcaacgggtc 307 Gly Ala Tyr Val Lys Gly Ala Ile Glu Tyr Phe Thr Lys Gln Asn Phe Tyr Ala Thr Gly
1021 atggcgttctctcaggtgagaactatgtttatcaggccaatgcagctgataactactatg 327 His Gly Val Leu Ser Gly Glu Asn Tyr Val Tyr Gin Ala Asn Ala Ala Asp Asn Tyr Tyr
1081 ccgtcaagagtgatggcacaagcttgagaatgtggtggcacaacaaccttggaggcggtc 347 Ala Val Lys Ser Asp Gly Thr Ser Leu Arg Met Trp Trp His Asn Asn Leu Gly Gly Gly
1141 aaacatggttttgcatggggcccaccattaatgcaccgccgtttaatacgatggacttca 367 Gln Thr Trp Phe Cys Met Gly Pro Thr Ile Asn Ala Pro Pro Phe Asn Thr Met Asp Phe
1201 acggaaactctaatatttccagccggattagtgactataagcaggttggcgcttattttt 387 Asn Gly Asn Ser Asn Ile Ser Ser Arg Ile Ser Asp Tyr Lys Gln Val Gly Ala Tyr Phe
1261 tecaaacagacggaccggagatetacgaggacagtgttgtecatgacgtettetggcatg 407 Phe Gln Thr Asp Gly Pro Glu Ile Tyr Glu Asp Ser Val Val His Asp Val Phe Trp His
1321 ttaatgatgatgccatcaagacatattattccggagcttcaatttcacgagcaaccatct

2/6

- 427 Val Asn Asp Asp Ala Ile Lys Thr Tyr Tyr Ser Gly Ala Ser Ile Ser Arg Ala Thr Ile
- 1381 ggaagtgtcacaatgacccgatcatacagatgggctggacgtcacgaaatctcaccggaa
  447 Trp Lys Cys His Asn Asp Pro Ile Ile Gln Met Gly Trp Thr Ser Arg Asn Leu Thr Gly
- 1441 tcagcattgataacctgcacgtcatccacagagatatttcaaatctgaaacagtggttc
  467 lle Scr Ile Asp Asn Leu His Val Ile His Thr Arg Tyr Phe Lys Scr Glu Thr Val Val
- 1501 cttcagcaatcattggagcgtctccattctacgcaagtggaatgactgttgatcccagcg 487 Pro Ser Ala Ile Ile Gly Ala Ser Pro Phe Tyr Ala Ser Gly Met Thr Val Asp Pro Ser
- 1561 agtccatcagcatgaccatctctaacgtggtgtgtgtgagggtctatgcccctcactgttcc 507 Glu Scr Ile Scr Met Thr Ile Ser Asn Val Val Cys Glu Gly Leu Cys Pro Ser Leu Phe
- 1621 gtatcactccgcttcagagctacaacaaccttgttgtcaagaacgtggcctttcccgatg
  527 Arg Ile Thr Pro Leu Gln Ser Tyr Asn Asn Leu Val Val Lys Asn Val Ala Phe Pro Asp
- 1681 gactgcagacaaatccaatcggaataggagagagagtattataccagcagcttccggctgta 547 Gly Leu Gln Thr Asn Pro Ile Gly Ile Gly Glu Ser Ile Ile Pro Ala Ala Ser Gly Cys
- 1741 caatggacttggaaatcacaaactggaccgtcaaaggacaaaaagtcaccatgcaaaact
  567 Thr Met Asp Leu Glu Ile Thr Asn Trp Thr Val Lys Gly Gln Lys Val Thr Met Gln Asn
- 1801 ttcagtccgggtcacttggccagttcgatatcgatggttcatactggggtcaatggtcca 587 Phe Gln Ser Gly Ser Leu Gly Gln Phe Asp Ile Asp Gly Ser Tyr Trp Gly Gln Trp Ser
- 1861 taaac<u>taa</u>agctattcccattcacctgagtattttcgtgggttcaatgagttcttgttac 607 lle Asn \*
- 1921 tgatggggcccttgctagtggtaaaagtagagggacttgtcctcgccgggcgccaaggaa
- 2041 aaaaaaaaaaaa 2052

FIG. 1b

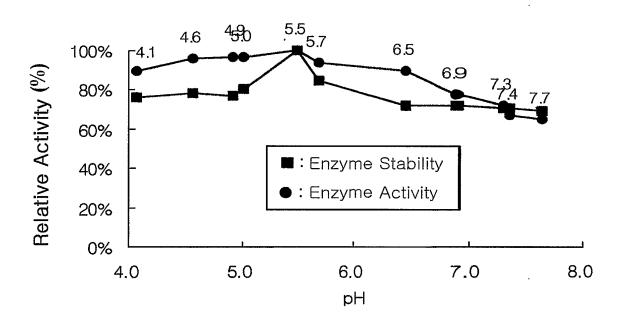


FIG. 2

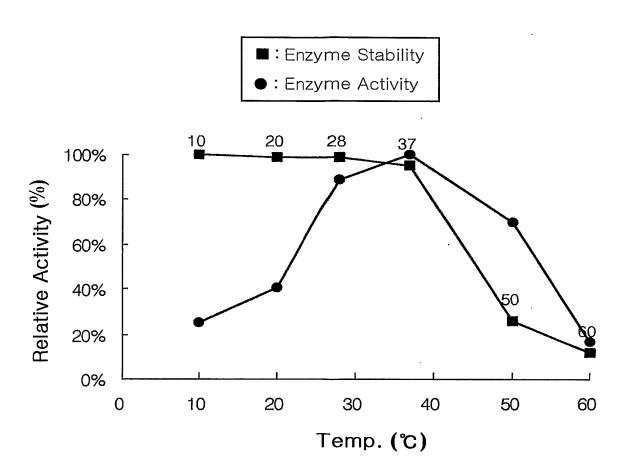


FIG. 3

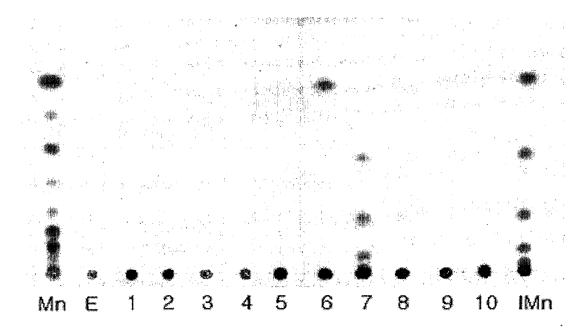


FIG. 4

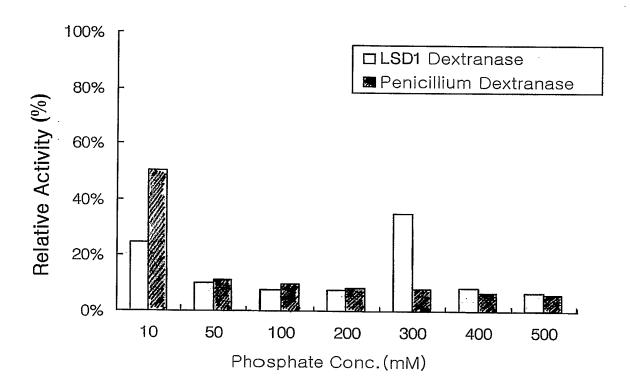


FIG. 5